



# STIC Search Report

## Biotech-Chem Library

STIC Database Tracking Number: 129546

TO: Leon Y Lum  
Location: REM/3D78/3C70  
Art Unit: 1641  
Sunday, August 15, 2004  
Case Serial Number: 10/044708

From: Paul Schulwitz  
Location: Biotech-Chem Library  
REM-1A65  
Phone: (571)272-2527

[paul.schulwitz@uspto.gov](mailto:paul.schulwitz@uspto.gov)

### Search Notes

Examiner Lum,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Paul Schulwitz  
Technical Information Specialist  
STIC Biotech/Chem Library  
(571)272-2527



## SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name: Leon Lum Examiner #: 80278 Date: 8/9/04  
 Art Unit: 1641 Phone Number 302-2878 Serial Number: 10/044708  
 Mail Box and Bldg/Room Location: Remsen Bldg. Results Format Preferred (circle): PAPER DISK E-MAIL  
Mailbox: 3070  
Room: 3078

If more than one search is submitted, please prioritize searches in order of need.

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Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc. if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of Invention: Isotope-coded ionization-enhancing reagents (ICIER) for high-throughput protein identification and quantitation using matrix-assisted laser desorption/ionization mass spectrometry

Inventors (please provide full names): Xiong-chang Qiu, Jack Wang, Rodney M. Hewick, Jack H. Wang

Earliest Priority Filing Date: 10/23/2000

\*For Sequence Searches Only\* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.

Please search the structures of Claim 9 (see attached) and the following search terms:  
 $\Delta$  = Deuterium

Matrix Assisted Laser Desorption/Ionization (MALDI)

Mass Spectrometry (MS)

MALDI-MS

Isotope Coded Affinity Tags (ICAT)

STAFF USE ONLY		Type of Search	Vendors and cost where applicable
Searcher: _____	NA Sequence (#) _____	STN <u>730.42</u>	
Searcher Phone #: _____	AA Sequence (#) _____	Dialog _____	
Searcher Location: _____	Structure (#) <u>2</u>	Questel/Orbit _____	
Date Searcher Picked Up: _____	Bibliographic _____	Dr.Link _____	
Date Completed: <u>8/5</u>	Litigation _____	Lexis/Nexis _____	
Searcher Prep & Review Time: <u>20</u>	Fulltext _____	Sequence Systems _____	
Clerical Prep Time: <u>100</u>	Patent Family _____	WWW/Internet _____	
Online Time: <u>25</u>	Other _____	Other (specify) _____	

WHAT IS CLAIMED IS:

1. A method for enhancing identification and relative quantitation of proteins and peptides using mass spectrometry (MS), said method comprising the steps of:

(a) reducing the disulfide bonds of a first sample from a biological mixture containing proteins and peptides;

(b) labeling proteins and peptides in the first sample with a reagent which comprises a thiol-specific reactive group attached to a guanidino group via a linker which can be differentially labeled;

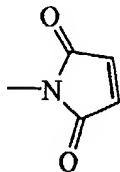
(c) separating the proteins and peptides from the sample;

(d) digesting the proteins to provide a mixture containing digestion peptides and peptides from the first sample; and

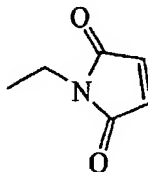
(e) subjecting the peptides of (d) to quantitative MS analysis and protein identification.

2. The method according to claim 1, wherein the peptides of (d) are subjected to matrix-assisted laser desorption/ionization (MALDI) - MS.

3. The method according to claim 1, wherein the reagent comprises a thiol-specific reactive group is selected from the group consisting of  $\alpha$ -haloacetyl ( $-X-CH_2CO-$ ,  $X = I, Br, \text{ or } Cl$ ) or a maleimide group having a structure selected from the group consisting of:



and



12.29 - 12.54

(25)

4. The method according to claim 1, wherein the linker comprises an alkyl chain having three to eight carbon atoms, optionally substituted with one or more amido groups, carboxy groups, or amino groups.

5. The method according to claim 1, wherein the proteins and peptides are further subjected to peptide mass mapping, said method further comprising the steps of:

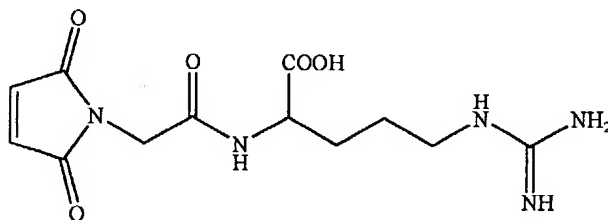
labeling proteins and peptides in a second sample with said reagent having heavy stable isotopes; and  
mixing the first and second samples prior to the separation step, wherein the reagent in the labeling step contains light stable isotopes.

6. The method according to claim 1, wherein the linker in the reagent of step (b) contains a substitution of four to twelve atoms with a stable isotope.

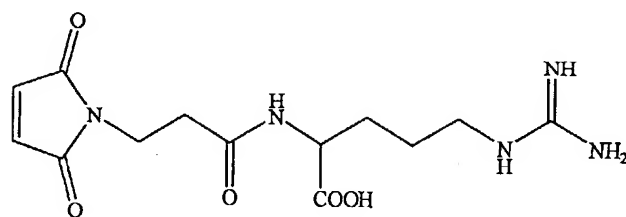
7. The method according to claim 6, wherein the linker contains seven stable isotopes.

8. The method according to claim 6, wherein the hydrogen atoms are substituted with deuterium.

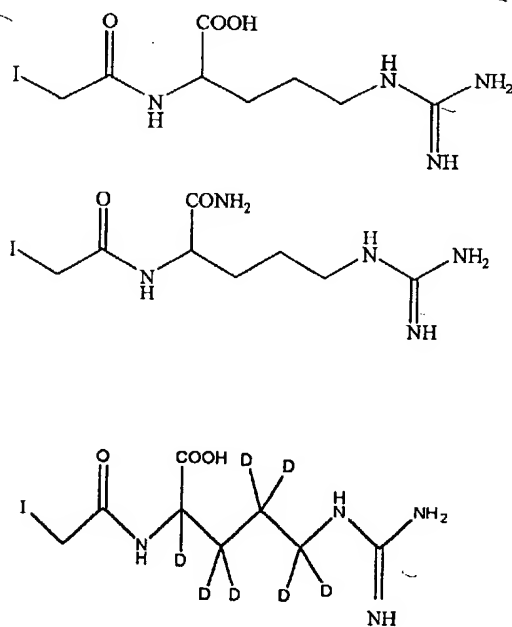
9. The method according to claim 5, wherein the reagent is selected from the group consisting of:



(LS)



and



10. The method according to claim 5, wherein the separation step is performed using one dimensional or two dimensional polyacrylamide gel electrophoresis (1D or 2D-PAGE), or liquid chromatography.

11. The method according to claim 1, wherein the digestion step is performed in-gel or in solution.

12. A method for preparing peptides for MALDI-MS and subsequent data analysis, said method comprising the steps of:

(a) reducing the disulfide bonds of proteins from biological samples;

(b) labeling proteins in one sample with a reagent which comprises a thiol-specific reactive group attached to a guanidino group via a linker which is differentially labeled with light stable isotopes;

(c) labeling proteins in a second sample with a reagent having heavy stable isotopes;

(d) mixing the first and second labeled samples;

(e) separating the proteins from the mixture;

(f) digesting the proteins, thereby providing peptides ready for MALDI-MS analysis and protein identification.

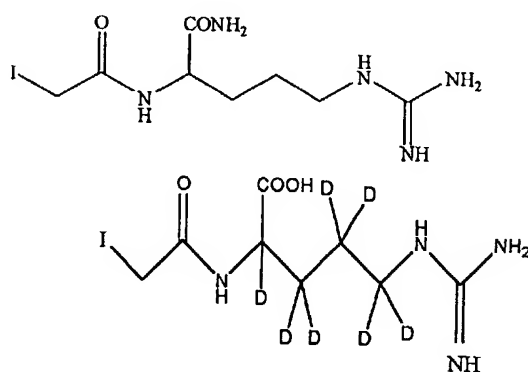
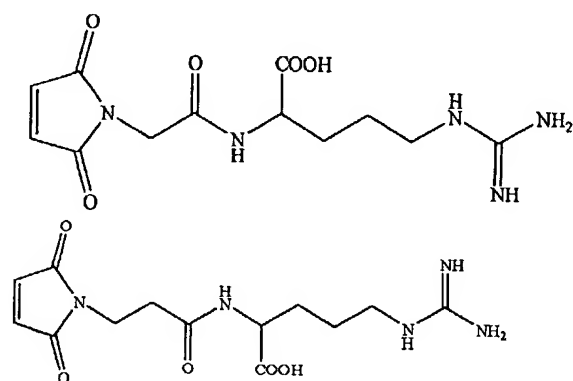
13. The method according to claim 11, wherein the digestion step is performed using trypsin.

14. A compound useful in quantitative analysis of protein mixtures, said compound comprising a thiol-specific reactive group attached to a guanidino group via a linker which can be differentially labeled with stable isotopes.

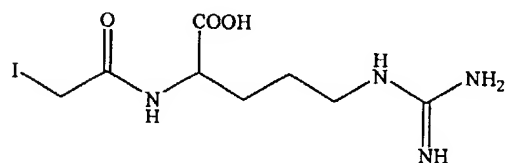
15. The compound according to claim 14, wherein the linker contains four to twelve stable isotopes.

16. The compound according to claim 14, wherein the linker contains a substitution of at least six hydrogen atoms with deuterium.

17. The compound according to claim 14, selected from the group consisting of:



and



18. A reagent kit for the analysis of proteins by mass spectrometric analysis that comprises a compound of claim 14 or claim 17.

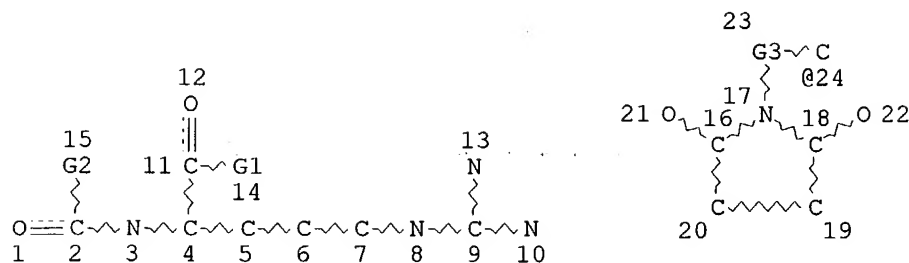
19. The reagent kit according to claim 18, comprising a set of substantially identical differentially labeled alkylating reagents.

20. The reagent kit according to claim 18, further comprising one or more proteolytic enzymes for use in digestion of proteins modified by said compounds.



L13

STR



I ~ C  
25 @26

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VAR G1=OH/NH2
VAR G2=24/26
REP G3=(0-4) C
NODE ATTRIBUTES:
DEFAULT MLEVEL IS ATOM
DEFAULT ELEVEL IS LIMITED
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GRAPH ATTRIBUTES:  
RING(S) ARE ISOLATED OR EMBEDDED  
NUMBER OF NODES IS 26

STEREO ATTRIBUTES: NONE

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L16 ANSWER 1 OF 1 HCAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 1968:78612 HCAPLUS

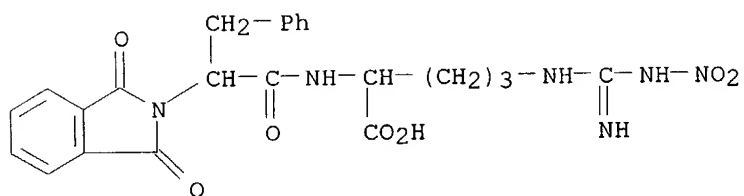
DOCUMENT NUMBER: 68:78612

TITLE: Potential antiviral agents. Carbobenzoxy di- and tripeptides active against measles and herpes viruses

AUTHOR(S): Nicolaides, Ernest D.; De Wald, Horace A.; Westland, Roger D.; Lipnik, Marilyn; Posler, Jeanette

CORPORATE SOURCE: Parke, Davis and Co., Ann Arbor, MI, USA

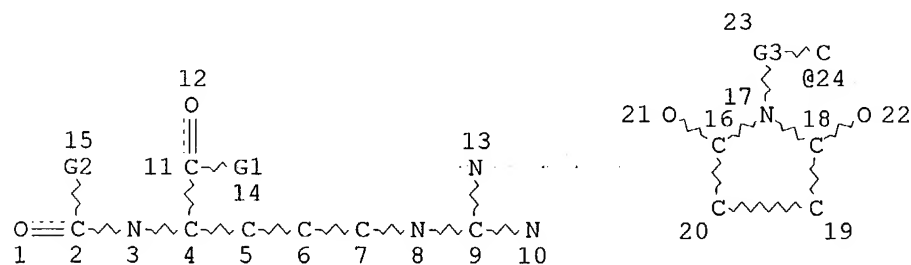
SOURCE: Journal of Medicinal Chemistry (1967), 11(1), 74-9  
CODEN: JMCMAR; ISSN: 0022-2623  
DOCUMENT TYPE: Journal  
LANGUAGE: English  
AB A large number of carbobenzoxy dipeptides, several tripeptides, and a number of alkyl, cycloalkyl, aryl, and heterocyclic amide derivs. of carbobenzoxy-L-and D-phenylalanine were synthesized. Many of the peptides were active against measles and herpes viruses.  
IT **17461-57-3P**  
RL: SPN (Synthetic preparation); PREP (Preparation)  
(preparation of)  
RN 17461-57-3 HCAPLUS  
CN Ornithine, N5-(nitroamidino)-N2-(L- $\alpha$ -phthalimidohydrocinnamoyl)-, L-  
(8CI) (CA INDEX NAME)



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L13

STR



I~C  
25 @26

VAR G1=OH/NH2

VAR G2=24/26

REP G3=(0-4) C

NODE ATTRIBUTES:

DEFAULT MLEVEL IS ATOM

DEFAULT ECLEVEL IS LIMITED

GRAPH ATTRIBUTES:

RING(S) ARE ISOLATED OR EMBEDDED

NUMBER OF NODES IS 26

STEREO ATTRIBUTES: NONE

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L20 1 SEA FILE=MARPAT ABB=ON PLU=ON L19/COM

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L20 ANSWER 1 OF 1 MARPAT COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 128:34581 MARPAT

TITLE: Preparation of acetylene derivatives for inhibition of matrix metalloproteases

INVENTOR(S): Dixon, Brian R.; Chen, Jinshan

PATENT ASSIGNEE(S): Bayer Corporation, USA; Dixon, Brian R.; Chen, Jinshan

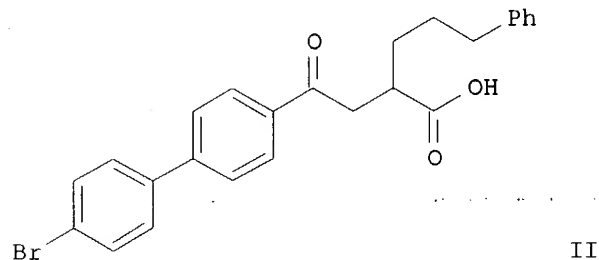
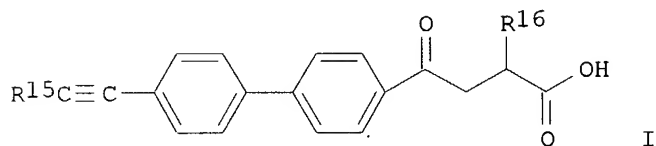
SOURCE: PCT Int. Appl., 71 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 2  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9743245	A1	19971120	WO 1997-US7921	19970512
W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
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ZA 9704031	A	19980219	ZA 1997-4031	19970509
HR 970245	B1	20020630	HR 1997-970245	19970509
AU 9729386	A1	19971205	AU 1997-29386	19970512
AU 710759	B2	19990930		
EP 912496	A1	19990506	EP 1997-923622	19970512
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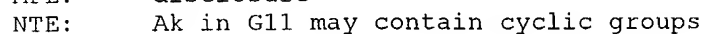
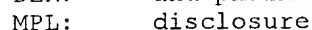
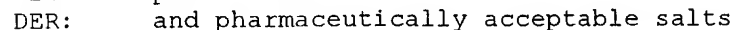
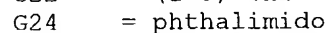
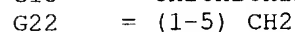
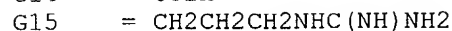
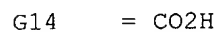
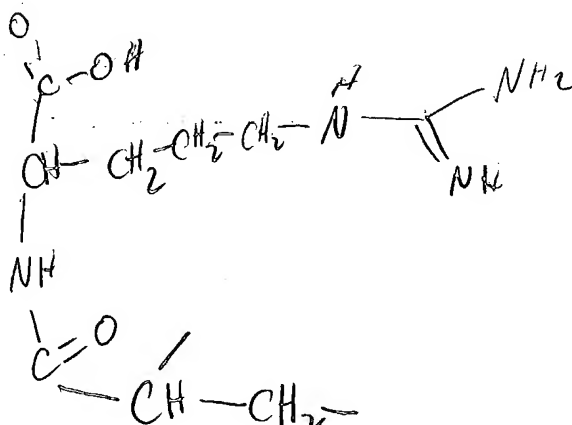
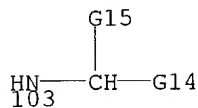
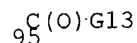
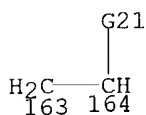
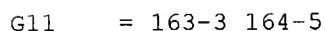
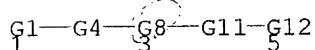
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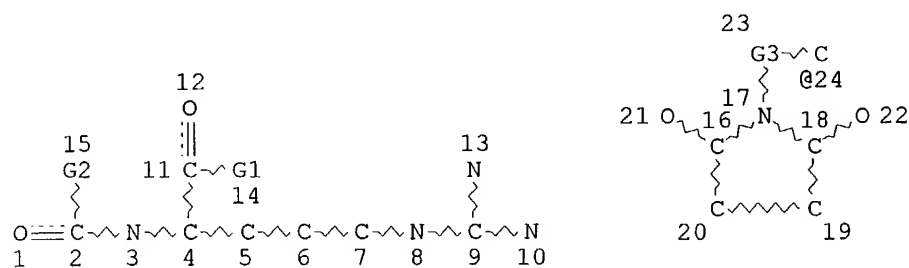
AB The title compds. [I; R15 = HOCH<sub>2</sub>, MeOCH<sub>2</sub>, CH<sub>3</sub>CO<sub>2</sub>CH<sub>2</sub>, EtOCO<sub>2</sub>CH<sub>2</sub>, HO(CH<sub>2</sub>)<sub>2</sub>, CH<sub>3</sub>CO<sub>2</sub>(CH<sub>2</sub>)<sub>2</sub>, HO<sub>2</sub>C(CH<sub>2</sub>)<sub>2</sub>, OHC(CH<sub>2</sub>)<sub>3</sub>, HO(CH<sub>2</sub>)<sub>4</sub>, Ph, etc.; R16 = Ph(CH<sub>2</sub>)<sub>3</sub>, phthalimidoethyl] are prepared I are useful for inhibiting matrix

metalloproteases and, therefore, combating conditions to which MMP's contribute, such as osteoarthritis, rheumatoid arthritis, septic arthritis, periodontal disease, corneal ulceration, proteinuria, aneurysmal aortic disease, dystrophic epidermolysis, bullosa, conditions leading to inflammatory responses, osteopenias mediated by MMP activity, temporomandibular joint disease, demyelating diseases of the nervous system, tumor metastasis or degenerative cartilage loss following traumatic joint injury, and coronary thrombosis from atherosclerotic plate rupture. Thus, compound (II) was reacted with HOCH<sub>2</sub>C.tplbond.CH in the presence of Et<sub>2</sub>NH, CuI, and trans-dichlorobis(triphenylphosphine)palladate to give I [R15 = HOCH<sub>2</sub>, R16 = Ph(CH<sub>2</sub>)<sub>3</sub>], which showed IC<sub>50</sub> of 21 μM against MMP-3.

## MSTR 2



STR



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VAR G1=OH/NH2
VAR G2=24/26
REP G3=(0-4) C
NODE ATTRIBUTES:
DEFAULT MLEVEL IS ATOM
DEFAULT ECLEVEL IS LIMITED

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STEREO ATTRIBUTES: NONE
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